

RAW SEQUENCE LISTING
PATENT APPLICATION: US/08/962,560B

DATE: 09/14/2000
TIME: 17:33:46

Input Set : A:\PTO.txt
Output Set: N:\CRF3\09142000\H962560B.raw

Does Not Comply
Corrected Diskette Needed

1, 2, 3
See pp.

4 <110> APPLICANT: Hilton, Douglas J.
5 Alexander, Warren S.
6 Viney, Elizabeth M.
7 Wilson, Tracy A.
8 Richardson, Rachel
9 Starr, Robyn
10 Nicholson, Sandra E.
11 Metcalf, Donald
12 Nicola, Nicos A.
14 <120> TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC AGENTS
16 <130> FILE REFERENCE: Davies Collision Cave
18 <140> CURRENT APPLICATION NUMBER: 08/962,560B
19 <141> CURRENT FILING DATE: 1997-10-31
21 <160> NUMBER OF SEQ ID NOS: 68
23 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

137 <210> SEQ ID NO: 4
138 <211> LENGTH: 212
139 <212> TYPE: PRT
140 <213> ORGANISM: Mus musculus
142 <400> SEQUENCE: 4
143 Met Val Ala Arg Asn Gln Val Ala Ala Asp Asn Ala Ile Ser Pro Ala
144 1 5 10 15
146 Ala Glu Pro Arg Arg Arg Ser Glu Pro Ser Ser Ser Ser Ser Ser
147 20 25 30
149 Ser Pro Ala Ala Pro Val Arg Pro Arg Pro Cys Pro Ala Val Pro Ala
150 35 40 45
152 Pro Ala Pro Gly Asp Thr His Phe Arg Thr Phe Arg Ser His Ser Asp
153 50 55 60
155 Tyr Arg Arg Ile Thr Arg Thr Ser Ala Leu Leu Asp Ala Cys Gly Phe
156 65 70 75 80
158 Tyr Trp Gly Pro Leu Ser Val His Gly Ala His Glu Arg Leu Arg Ala
159 85 90 95
E--> 161 Glu Pro Va Gly Thr Phe Leu Val Arg Asp Ser Arg Gln Arg Asn Cys
162 100 105 110
164 Phe Phe Ala Leu Ser Val Lys Met Ala Ser Gly Pro Thr Ser Ile Arg
165 115 120 125
167 Val His Phe Gln Ala Gly Arg Phe His Leu Asp Gly Ser Arg Glu Thr
168 130 135 140
171 Phe Asp Cys Leu Phe Glu Leu Leu Glu His Tyr Val Ala Ala Pro Arg
172 145 150 155 160
174 Arg Met Leu Gly Ala Pro Leu Arg Gln Arg Arg Val Arg Pro Leu Gln
175 165 170 175
177 Glu Leu Cys Arg Gln Arg Ile Val Ala Ala Val Gly Arg Glu Asn Leu

Invalid amino acid
designator.

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178 180 185 190
180 Ala Arg Ile Pro Leu Asn Pro Val Leu Arg Asp Tyr Leu Ser Ser Phe
181 195 200 205
183 Pro Phe Gln Ile
184 210
1644 <210> SEQ ID NO: 24
1645 <211> LENGTH: 2019
1646 <212> TYPE: DNA
1647 <213> ORGANISM: Mus musculus
1649 <220> FEATURE:
1650 <221> NAME/KEY: UNSURE
1651 <222> LOCATION: (1981)
1652 <223> OTHER INFORMATION: N is unsure
1654 <220> FEATURE:
1655 <221> NAME/KEY: UNSURE
1656 <222> LOCATION: (1992)
1657 <223> OTHER INFORMATION: N is unsure
1659 <220> FEATURE:
1660 <221> NAME/KEY: UNSURE
1661 <222> LOCATION (2010) → n is at location 2000
1662 <223> OTHER INFORMATION: N is unsure
1664 <400> SEQUENCE: 24
1665 ggacagggc ggggtcaggc cggaggctga ggaccaagta ggcatggcgg agggcgggac 60
1667 cggcccccgcgat ggacggggccg cgcaggatcct aatctgaaagg agtggctgag 120
1669 ggacggatc tgggaccatc cactggacca ctgtgacgat acaagactcc atgatgcagc 180
1671 ctatgtatggg gacccatccaga cccctcaggaa cctactgca gaggagatc accggagcc 240
1673 catcaatgtatc aagtctgtttt ggtgctgccc ctggcttccc tgacacccatc tgaggatcgc 300
1675 agccactgtca ggcacatggga actgtgtggaa ctcccttcata cgcaaaagggg ccgagggtgaa 360
1677 cctgggtatgtcaaggggc agactgtccct gtatgtggat gtatgtggat ggcactgtga 420
1679 gaccaactgtatccatggg aagctggatc tgatccaaac ggcagccggc accaccgcag 480
1681 cactccgtgtt accatgtatc ytcgtgtggg tagggacgatc atccatgtatc 540
1683 gatggggca gatgttgcata tcaaccatc tctgtatccatc gacacccggc ccccttttc 600
1685 acggccgcata accctccatgg tggctgtcc tctatacatc atgtgtccctt accataactt 660
1687 teagtgttc accgtgtccat tgcaggatcggg gccaatccatc gacttcaattt gcaatggccc 720
1689 tgtaacacacc caggaggatc acaggatcggatc ccctgggtatgt gtcatggatg ctgtccgtc 780
1691 ccatgtgtgtt gaaaggcgcct tctgtgatct gttgttagat ttttgagccaa acctgtatc 840
1693 ggtgaatgtt gaaatccatgg gcccacgggc aagaggcaga agaaatgtt atccctggc 900
1695 ctgcaggatc tttaaaaggccatc ccagaatgtt tcccaaggacc ttgtgtatgtt tgccgggt 960
1697 ggctgtgaga agacgttccatc gcaatacccg actgtatccatc gttcccttcgc tgccgtgcc 1020
1699 agacccatata aagaatgtt tgcattatgtatc gtagcatccatc catgcgttc tgactgtatc 1080
1701 gtggaaaggccg atcacatgtca gttttttttttt gacacatccatc tggcatccatc ggaaccatgg 1140
1703 ccctgtgtgc cagcttgcatc ctggctgtc atgtatgtatc gttttttttttt gttttttttttt 1200
1705 actgtgtatccatc tttttttttttt gttttttttttt gttttttttttt gttttttttttt 1260
1707 gaggcacata caaaatccatc tttttttttttt tttttttttttt tttttttttttt 1320
1709 caatgtgtgc aacatgggc gggctgtgtgtt attggccatccatc tttttttttttt 1380
1711 aggtatgtatc tttttttttttt gttttttttttt gttttttttttt gttttttttttt 1440
E--> 1713 caaaacagaa agagggtttgtt taagaaaaaaa tttttttttttt gttttttttttt 1550 →
1715 actgtgtatccatc tttttttttttt tttttttttttt tttttttttttt tttttttttttt 1560
1717 ttctttttttttt tttttttttttt tttttttttttt tttttttttttt tttttttttttt 1620

miscount typo
should be 1500

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1719 gttctatctt attgcacaga ggtcccaaga cagtttcat agggcaccat ctgctctgcc 1680
1721 aagggttttc tgatgttta ccttgggat cttcagacag tggttaccc taggagacc 1740
1723 acctggact aaccattaaag tgactgccc cattcagatc agggaccatc ttaatagac 1800
1725 tcaactgcac agaagatgac acgggtgc tttcagaca ctccatata 1860
1727 ggaagtttga aatgttgg gtcacctgg ttgttccag gtcataactt cttgggttcc 1920
1729 cactaaracc agpatatcc tttttttgg gttgactgtt ccctccccac tttccttcaa 1980
W--> 1731 ncccaatgac cttttgtt n gttgcttcc ctaaaaktt 2019

n is at location 2000, not 2010

F.Y.I.

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 09/14/2000
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L:161 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:161 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:1085 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1713 M:254 E: No. of Bases conflict, LENGTH:Input:1550 Counted:1500 SEQ:24
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1916 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1920 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:2524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:2679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:3416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51